```
SEQUENCE LISTING
<110 >Tonen Corporation
<120 >Method for Detection or Measurement of Hepatitis C V
      irus
<160 >8
<210 >1
<211 >177
<212 >PRT
<213 >Hepatitis C virus
<400 >1
 Met Lys Ala Ile Phe Val Leu Lys Gly Ser Leu Asp Arg Asp Pro Glu
                                        10
 Phe Met Gly Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr
               20
                                    25
                                                        30
 Asn Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val
           35
                                40
 Gly Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg
                           55
 Ala Thr Arg Lys Thr Ser Lys Arg Ser Gln Pro Arg Gly Gly Arg Arg
                       70
  65
                                            75
                                                                 80
 Pro Ile Pro Lys Asp Arg Arg Ser Thr Gly Lys Ser Trp Gly Lys Pro
                   85
                                        90
 Gly Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly
                                   105
 Trp Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp
          115
                              120
                                                   125
 Pro Arg His Arg Ser Arg Asn Val Gly Lys Val Ile Asp Thr Leu Thr
      130
                          135
                                               140
 Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Phe Arg Val Gly Ala Phe
                      150
                                           155
 Leu Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu
                  165
                                       170
                                                           175
 Asp
```

<210 >2

```
<211 >160
<212 >TRP
<213 >Hepatitis C virus
<400 >2
 Met Gly Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
                                        10
 Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly
                                    25
               20
                                                         30
 Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
           35
                                40
                                                    45
 Thr Arg Lys Thr Ser Lys Arg Ser Gln Pro Arg Gly Gly Arg Arg Pro
                            55
                                                60
 Ile Pro Lys Asp Arg Arg Ser Thr Gly Lys Ser Trp Gly Lys Pro Gly
  65
                       70
                                            75
                                                                 80
 Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp
                   85
                                        90
                                                             95
 Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
                                   105
 Arg His Arg Ser Arg Asn Val Gly Lys Val Ile Asp Thr Leu Thr Cys
          115
                               120
                                                   125
 Gly Phe Ala Asp Leu Met Gly Tyr Ile Phe Arg Val Gly Ala Phe Leu
      130
                          135
                                               140
 Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
 145
                      150
                                           155
                                                                160
<210 >3
<211 >20
<212 >PRT
<213 >Artificial Sequence
<220 >
<223 >
<400 >3
 Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly Gly Val Tyr Leu
                                                             15
                                        10
```

```
Leu Pro Arg Arg
               20
<210 >4
<211 >10
<212 >PRT
<213 >Artificial Sequence
<220 >
<223 >
<400 >4
 Gly Pro Arg Leu Gly Val Arg Ala Thr Arg
                    5
                                        10
<210 >5
<211 >21
<212 >PRT
<213 >Artificial Sequence
<220 >
<223 >
<400 >5
  Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro Arg His Arg
    1
                    5
                                        10
                                                             15
 Ser Arg Asn Val Gly
               20
<210 >6
<211 >20
<212 >PRT
<213 >Artificial Sequence
<220 >
<230 >
<400 >6
 Asp Pro Arg His Arg Ser Arg Asn Val Gly Lys Val Lle Asp Thr Leu
    1
                                        10
                                                             15
 Thr Cys Gly Phe
               20
<210 >7
```

<211	>24	
<212	DNA	
<213	Artificial Sequence	
<220	Probe	
<230	Synthetic DNA	
<400	>7	
gaa	ttcatgg gcacgaatcc taaa	24
<210	>8	
<211	>21	
<212	DNA	
<213	Artificial Sequence	
<220	>Probe	
<230	Synthetic DNA	
<400	8	
tta	dtcctcc agaacccgga c	21